

Sequence alignment of 14 proteins (39976873.y, 32411113.y, 6323394.y, 19112506.y, AK110115.q, 6322366.y, 1f3m_Paka, 39973941.y, 32421133.y, 19113917.y, 39976873.y, 32411113.y, 6323394.y, 19112506.y, AK110115.q, 6322366.y, 1f3m_Paka, 39973941.y, 32421133.y, 19113917.y) across 1800 amino acids. The alignment highlights conservation of residues across the sequence. A color scale at the top indicates hydrophobicity: blue (most hydrophobic), green, yellow, orange, red (most hydrophilic).